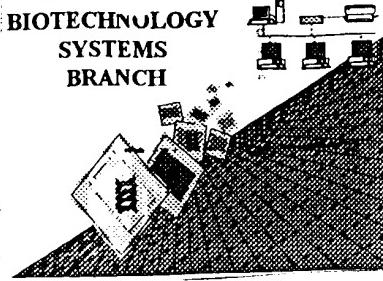


RAW SEQUENCE LISTING

ERROR REPORT



04C0
3-7-01

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/727,855

Source: OIPE

Date Processed by STIC: 12/14/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
 - 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
- FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§ 1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/722,855

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- | | |
|--|--|
| 1 <input type="checkbox"/> Wrapped Nucleic | The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping". |
| 2 <input type="checkbox"/> Wrapped Aminos | The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping". |
| 3 <input type="checkbox"/> Incorrect Line Length | The rules require that a line not exceed 72 characters in length. This includes spaces. |
| 4 <input type="checkbox"/> Misaligned Amino Acid Numbering | The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers. |
| 5 <input type="checkbox"/> Non-ASCII | This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed. |
| 6 <input type="checkbox"/> Variable Length | Sequence(s) <input type="checkbox"/> contain n's or Xaa's which represent more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing. |
| 7 <input type="checkbox"/> PatentIn ver. 2.0 "bug" | A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) <input type="checkbox"/> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences. |
| 8 <input type="checkbox"/> Skipped Sequences (OLD RULES) | Sequence(s) <input type="checkbox"/> missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s). |
| 9 <input type="checkbox"/> Skipped Sequences (NEW RULES) | Sequence(s) <input type="checkbox"/> missing. If intentional, please use the following format for each skipped sequence:
<210> sequence id number
<400> sequence id number
000 |
| 10 <input checked="" type="checkbox"/> Use of n's or Xaa's (NEW RULES) | Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents. |
| 11 <input type="checkbox"/> Use of <213>Organism (NEW RULES) | Sequence(s) <input type="checkbox"/> are missing this mandatory field or its response. |
| 12 <input type="checkbox"/> Use of <220>Feature (NEW RULES) | Sequence(s) <input type="checkbox"/> are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules) |
| 13 <input type="checkbox"/> PatentIn ver. 2.0 "bug" | Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk. |

Report Date: 10/10/2012
Report Type: Final

Report Status: Complete
Report Source: Internal Audit

fr 6

CK ~~Review and Implement the Policy, Report the findings and
recommendations to the Board of Directors.~~

1.0 Executive Summary

1.1 Purpose of the Audit

The purpose of this audit was to review and evaluate the

internal control system of the organization.

The audit was conducted to determine if the organization's

internal control system is effective and efficient in carrying

out its mission and objectives.

The audit was also conducted to identify any weaknesses in

the internal control system and to recommend corrective

actions to be taken by the organization.

The audit was conducted in accordance with generally accepted

auditing standards and procedures.

The audit was conducted by a team of auditors from the

internal audit department of the organization.

The audit was conducted to ensure that the organization's

internal control system is effective and efficient in carrying

out its mission and objectives.

The audit was conducted to identify any weaknesses in

the internal control system and to recommend corrective

actions to be taken by the organization.

The audit was conducted in accordance with generally accepted

auditing standards and procedures.

The audit was conducted by a team of auditors from the

internal audit department of the organization.

The audit was conducted to ensure that the organization's

internal control system is effective and efficient in carrying

out its mission and objectives.

The audit was conducted to identify any weaknesses in

the internal control system and to recommend corrective

actions to be taken by the organization.

The audit was conducted in accordance with generally accepted

auditing standards and procedures.

The audit was conducted by a team of auditors from the

internal audit department of the organization.

and the following day, the first two were found to have died. The third was still alive but had lost its right wing. It was released and was seen to fly away. The fourth bird was captured again on 20th April and was found to have lost its left wing. It was released and was seen to fly away.

The fifth bird was captured on 21st April and was found to have lost its right wing. It was released and was seen to fly away.

The sixth bird was captured on 22nd April and was found to have lost its left wing. It was released and was seen to fly away.

The seventh bird was captured on 23rd April and was found to have lost its right wing. It was released and was seen to fly away.

The eighth bird was captured on 24th April and was found to have lost its left wing. It was released and was seen to fly away.

The ninth bird was captured on 25th April and was found to have lost its right wing. It was released and was seen to fly away.

The tenth bird was captured on 26th April and was found to have lost its left wing. It was released and was seen to fly away.

The eleventh bird was captured on 27th April and was found to have lost its right wing. It was released and was seen to fly away.

The twelfth bird was captured on 28th April and was found to have lost its left wing. It was released and was seen to fly away.

The thirteenth bird was captured on 29th April and was found to have lost its right wing. It was released and was seen to fly away.

The fourteenth bird was captured on 30th April and was found to have lost its left wing. It was released and was seen to fly away.

The fifteenth bird was captured on 1st May and was found to have lost its right wing. It was released and was seen to fly away.

The sixteenth bird was captured on 2nd May and was found to have lost its left wing. It was released and was seen to fly away.

The seventeenth bird was captured on 3rd May and was found to have lost its right wing. It was released and was seen to fly away.

The eighteenth bird was captured on 4th May and was found to have lost its left wing. It was released and was seen to fly away.

The nineteenth bird was captured on 5th May and was found to have lost its right wing. It was released and was seen to fly away.

The twentieth bird was captured on 6th May and was found to have lost its left wing. It was released and was seen to fly away.

Digitized by Google

$\{ \omega_i \}_{i=1}^n$ (1)

1996-1997 学年第一学期期中考试

2010-07-01 00:00:00 UTC

1214 (11)

4/27/85

6

<210> 10
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sod1 (sense
primer for cloning of SOD genes)

<400> 10
aarcaycayc araqntaygt naa

23

see item 10 on Error Summary Sheet

<210> 11
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Sod4 (antisense
primer for cloning of SOD genes)

<400> 11
gcccanccng a[n]ccytgnac [n]cc

23

see item 10

FYI:

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

